

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	18	Ruvkun NEAR Gary	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2005/07/11 08:55
L2	39	DAF-16	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2005/07/11 08:55
L3	7323	ELEGANS	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2005/07/11 08:56
L4	538	AFx fchr	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L5	14	L2 and L3 and L4	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L6	65867	insulin	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L7	119	L6 and L4	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L8	41	L7 and L3	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L9	29	(AFx FKHR def-16).clm.	US-PGPUB; USPAT; EPO; DERWENT	OR	ON	2005/07/11 08:56
L10	48	(glucose insulin) AND (afx OR fchr OR def-16)	US-PGPUB; USPAT; EPO; DERWENT	SAME	ON	2005/07/11 08:56

=> d his

(FILE 'HOME' ENTERED AT 08:59:05 ON 11 JUL 2005)

FILE 'MEDLINE, CANCERLIT, AGRICOLA, CAPLUS, SCISEARCH' ENTERED AT
09:02:32 ON 11 JUL 2005

L1 461 S DAF-16
L2 1536 S AFX OR FKHR
L3 19693 S C. ELEGAN?
L4 21 S L1 (L) L2 (L) L3
L5 7 DUP REM L4 (14 DUPLICATES REMOVED)
L6 0 S L5 AND PY<=1997
E RUVKUN GARY?/AU

=> d an ti so au ab l5 1-7

L5 ANSWER 1 OF 7 MEDLINE on STN DUPLICATE 1
AN 2003538461 MEDLINE
TI Convergence of peroxisome proliferator-activated receptor gamma and Foxo1
signaling pathways.
SO Journal of biological chemistry, (2003 Nov 14) 278 (46) 45485-91.
Electronic Publication: 2003-09-09.
Journal code: 2985121R. ISSN: 0021-9258.
AU Dowell Paul; Otto Tamara C; Adi Saleh; Lane M Daniel
AB The forkhead factor Foxo1 (or **FKHR**) was identified in a yeast
two-hybrid screen as a peroxisome proliferator-activated receptor (PPAR)
gamma-interacting protein. Foxo1 antagonized PPARGamma activity and vice
versa indicating that these transcription factors functionally interact in
a reciprocal antagonistic manner. One mechanism by which Foxo1
antagonizes PPARGamma activity is through disruption of DNA binding as
Foxo1 inhibited the DNA binding activity of a PPARGamma/retinoid X
receptor alpha heterodimeric complex. The *Caenorhabditis elegans* nuclear
hormone receptor, DAF-12, interacted with the *C. elegans*
forkhead factor, **DAF-16**, paralleling the interaction
between PPARGamma and Foxo1. *daf-12* and *daf-16* have
been implicated in *C. elegans* insulin-like signaling
pathways, and PPARGamma and Foxo1 likewise have been linked to mammalian
insulin signaling pathways. These results suggest a convergence of
PPARGamma and Foxo1 signaling that may play a role in insulin action and
the insulinomimetic properties of PPARGamma ligands. A more general
convergence of nuclear hormone receptor and forkhead factor pathways may
be important for multiple biological processes and this convergence may be
evolutionarily conserved.

L5 ANSWER 2 OF 7 MEDLINE on STN DUPLICATE 2
AN 2002664265 MEDLINE
TI Effects of aging and caloric restriction on the gene expression of Foxo1,
3, and 4 (**FKHR**, **FKHRL1**, and **AFX**) in the rat skeletal muscles.
SO Microscopy research and technique, (2002 Nov 15) 59 (4) 331-4.
Journal code: 9203012. ISSN: 1059-910X.
AU Furuyama Tatsuo; Yamashita Hitoshi; Kitayama Kazuko; Higami Yoshikazu;
Shimokawa Isao; Mori Nozomu
AB In *C. elegans*, insulin-like hormone signal pathway
plays a significant role in longevity. In particular, *daf-16*
gene product is indispensable factor for this
lifespan-extension. This signal pathway is critical for dauer formation,
which is a similar state to hibernation in mammals. We examined the
expression level of mammalian *daf-16* homologues, Foxo
1,3, and 4 (**FKHR**, **FKHRL1**, and **AFX**) mRNAs in the rat
skeletal muscles during aging and in 30% caloric restricted of ad libitum
fed. The expression level of **AFX** mRNA was significantly higher
at 6 and 12 months than at 3 and 26 months, and **FKHRL1** expression was
significantly higher at 6 months than at 3 and 26 months but **FKHR**
expression showed no significant change with age. We observed a
characteristic expression of **AFX** and **FKHR** mRNAs to be
significantly higher in the second day in caloric restriction by
every-other-day feeding than in ad libitum fed. This suggests that
caloric restriction may increase the expression of **FKHR**-family
genes and prevent the aging process in the skeletal muscles.
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L5 ANSWER 3 OF 7 MEDLINE on STN DUPLICATE 3
 AN 2001308673 MEDLINE
 TI Phosphatidylinositol 3-kinase signaling inhibits DAF-16 DNA binding and function via 14-3-3-dependent and 14-3-3-independent pathways.
 SO Journal of biological chemistry, (2001 Apr 20) 276 (16) 13402-10.
 Electronic Publication: 2000-12-20.
 Journal code: 2985121R. ISSN: 0021-9258.
 AU Cahill C M; Tzivion G; Nasrin N; Ogg S; Dore J; Ruvkun G; Alexander-Bridges M
 AB In *Caenorhabditis elegans*, an insulin-like signaling pathway to phosphatidylinositol 3-kinase (PI 3-kinase) and AKT negatively regulates the activity of **DAF-16**, a Forkhead transcription factor. We show that in mammalian cells, **C. elegans DAF-16** is a direct target of AKT and that AKT phosphorylation generates 14-3-3 binding sites and regulates the nuclear/cytoplasmic distribution of **DAF-16** as previously shown for its mammalian homologs **FKHR** and **FKHRL1**. In vitro, interaction of AKT- phosphorylated **DAF-16** with 14-3-3 prevents **DAF-16** binding to its target site in the insulin-like growth factor binding protein-1 gene, the insulin response element. In HepG2 cells, insulin signaling to PI 3-kinase/AKT inhibits the ability of a GAL4 DNA binding domain/**DAF-16** fusion protein to activate transcription via the insulin-like growth factor binding protein-1-insulin response element, but not the GAL4 DNA binding site, which suggests that insulin inhibits the interaction of **DAF-16** with its cognate DNA site. Elimination of the **DAF-16/1433** association by mutation of the AKT/14-3-3 sites in **DAF-16**, prevents 14-3-3 inhibition of **DAF-16** DNA binding and insulin inhibition of **DAF-16** function. Similarly, inhibition of the **DAF-16/14-3-3** association by exposure of cells to the PI 3-kinase inhibitor LY294002, enhances **DAF-16** DNA binding and transcription activity. Surprisingly constitutively nuclear **DAF-16** mutants that lack AKT/14-3-3 binding sites also show enhanced DNA binding and transcription activity in response to LY294002, pointing to a 14-3-3-independent mode of regulation. Thus, our results demonstrate at least two mechanisms, one 14-3-3-dependent and the other 14-3-3-independent, whereby PI 3-kinase signaling regulates **DAF-16** DNA binding and transcription function.

L5 ANSWER 4 OF 7 MEDLINE on STN DUPLICATE 4
 AN 2001699838 MEDLINE
 TI Regulation of *C. elegans* DAF-16 and its human ortholog FKHRL1 by the daf-2 insulin-like signaling pathway.
 SO Current biology : CB, (2001 Dec 11) 11 (24) 1950-7.
 Journal code: 9107782. ISSN: 0960-9822.
 AU Lee R Y; Hench J; Ruvkun G
 AB *C. elegans* insulin-like signaling regulates metabolism, development, and life span. This signaling pathway negatively regulates the activity of the forkhead transcription factor **DAF-16**. **daf-16** encodes multiple isoforms that are expressed in distinct tissue types and are probable orthologs of human **FKHRL1**, **FKHR**, and **AFX**. We show that human **FKHRL1** can partially replace **DAF-16**, proving the orthology. In mammalian cells, insulin and insulin-like growth factor signaling activate AKT/PKB kinase to negatively regulate the nuclear localization of **DAF-16** homologs (reviewed in). We show that the absence of AKT consensus sites on **DAF-16** is sufficient to cause dauer arrest in **daf-2(+)** animals, proving that **daf-16** is the major output of insulin signaling in *C. elegans*. **FKHR**, **FKHRL1**, and **AFX** may similarly be the major outputs of mammalian insulin signaling. **daf-2** insulin signaling, via AKT kinases, negatively regulates **DAF-16** by controlling its nuclear localization. Surprisingly, we find that **daf-7** TGF-beta signaling also regulates **DAF-16** nuclear localization specifically at the time when the animal makes the commitment between diapause and reproductive development. **daf-16** function is supported by the combined action of two distinct promoter/enhancer elements, whereas the coding sequences of two major

DAF-16 isoforms are interchangeable. Together, these observations suggest that the combined effects of transcriptional and posttranslational regulation of **daf-16** transduce insulin-like signals in *C. elegans* and perhaps more generally.